SEQUENCE LISTING

<110> Bristol-Myers Squibb Company

<120> A NOVEL HUMAN LEUCINE-RICH REPEAT CONTAINING PROTEIN EXPRESSED PREDOMINATELY IN SMALL INTESTINE, HLRRS11

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gac gg Asp Gl	c gcg y Ala 15	gac Asp	gag Glu	ctg Leu	ccg Pro	gcg Ala 20	ctg Leu	ggg Gly	ggc Gly	ccc Pro	gag Glu 25	gcc Ala	gcg Ala	ccc Pro	158
tgc aca Cys Th: 30	a gac r Asp	ccc Pro	ttc Phe	gag Glu	gcg Ala 35	gcg Ala	agc Ser	ggc Gly	gcg Ala	cgg Arg 40	gtg Val	cta Leu	ggc Gly	GJÀ ààà	206
ctg cto Leu Leo 45	g agt ı Ser	aag Lys	gcg Ala	ctg Leu 50	ctg Leu	ccc Pro	acg Thr	gcc Ala	ctc Leu 55	ctg Leu	ctg Leu	gtg Val	acc Thr	acg Thr 60	254
cgc gcc Arg Ala	c gcc a Ala	gcc Ala	ccc Pro 65	ggg Gly	agg Arg	ctg Leu	cag Gln	ggc Gly 70	cgc Arg	ctg Leu	tgt Cys	tcc Ser	ccg Pro 75	cag Gln	302
tgc gcc Cys Ala	gag Glu	gtg Val 80	cgc Arg	ggc Gly	ttc Phe	tcc Ser	gac Asp 85	aag Lys	gac Asp	aag Lys	aag Lys	aag Lys 90	tat Tyr	ttc Phe	350
tac aaq Tyr Lys	ttc Phe 95	ttc Phe	cgg Arg	gat Asp	gag Glu	agg Arg 100	agg Arg	gcc Ala	gag Glu	cgc Arg	gcc Ala 105	tac Tyr	cgc Arg	ttc Phe	398
gtg aag Val Lys 110	Glu	aac Asn	gag Glu	acg Thr	ctg Leu 115	ttc Phe	gcg Ala	ctg Leu	tgc Cys	ttc Phe 120	gtg Val	ccc Pro	ttc Phe	gtg Val	446
tgc tgg	atc	gtg	tgc	acc	gtg	ctg	cgc	cag	cag	ctg	gag	ctc	ggt	cgg	494

Cys 125	Trp	Ile	· Val	. Cys	Thr 130		Leu	Arg	Gln	Gln 135		Glü	ı Leu	Gly	Arg 140	
gac Asp	ctg Leu	tcg Ser	cgc Arg	acg Thr 145	Ser	: aag · Lys	acc Thr	acc Thr	acg Thr 150	Ser	gtg Val	tac Tyr	ctg Leu	ctt Leu 155	ttc Phe	542
atc Ile	acc Thr	agc Ser	gtt Val 160	Leu	ago Ser	tcg Ser	gct Ala	ccg Pro 165	Val	gcc Ala	gac Asp	ggg	ccc Pro 170	Arg	ttg Leu	590
cag Gln	ggc	gac Asp 175	Leu	cgc Arg	aat Asn	ctg Leu	tgc Cys 180	Arg	ctg Leu	gcc Ala	cgc Arg	gag Glu 185	Gly	gtc Val	ctc Leu	638
gga Gly	cgc Arg 190	agg Arg	gcg Ala	cag Gln	ttt Phe	gcc Ala 195	gag Glu	aag Lys	gaa Glu	ctg Leu	gag Glu 200	caa Gln	ctg Leu	gag Glu	ctt Leu	686
Arg 205	Gly	Ser	Lys	Val	Gln 210	Thr	Leu	ttt Phe	Leu	Ser 215	Lys	Lys	Glu	Leu	Pro 220	734
Gly	Val	Leu	Glu	Thr 225	Glu	Val	Thr	tac Tyr	Gln 230	Phe	Ile	Asp	Gln	Ser 235	Phe	782
GIn	Glu	Phe	Leu 240	Ala	Ala	Leu	Ser	tac Tyr 245	Leu	Leu	Glu	Asp	Gly 250	Gly	Val	830
Pro	Arg	Thr 255	Ala	Ala	Gly	Gly	Val 260	GJÀ aaa	Thr	Leu	Leu	Arg 265	Gly	Asp	Ala	878
Gln	Pro 270	His	Ser	His	Leu	Val 275	Leu	acc Thr	Thr	Arg	Phe 280	Leu	Phe	Gly	Leu	926
Leu 285	Ser	Ala	Glu	Arg	Met 290	Arg	Asp	atc Ile	Glu	Arg 295	His	Phe	Gly	Cys	Met 300	974
Val	Ser	Glu	Arg	Val 305	Lys	Gln	Glu	gcc Ala	Leu 310	Arg	Trp	Val	Gln	Gly 315	Gln	1022
GTÀ	GIn	Gly	Cys 320	Pro	Gly	Val	Ala	cca Pro 325	Glu	Val	Thr	Glu	Gly 330	Ala	Lys	1070
ggg Gly	ctc Leu	gag Glu 335	gac Asp	acc Thr	gaa Glu	gag Glu	cca Pro 340	gag Glu	gag Glu	gag Glu	gag Glu	gag Glu 345	gga Gly	gag Glu	gag Glu	1118
Pro	aac Asn 350	tac Tyr	cca Pro	ctg Leu	gag Glu	ttg Leu 355	ctg Leu	tac Tyr	tgc Cys	ctg Leu	tac Tyr 360	gag Glu	acg Thr	cag Gln	gag Glu	1166
gac Asp	gcg Ala	ttt Phe	gtg Val	cgc Arg	caa Gln	gcc Ala	ctg Leu	tgc Cys	cgg Arg	ttc Phe	ccg Pro	gag Glu	ctg Leu	gcg Ala	ctg Leu	1214

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Val Arg Cys C	ige eet get gga Cys Pro Ala Gly 100	cag gca ctg Gln Ala Leu 405	cgg ctg atc agc Arg Leu Ile Ser 410	tgc aga 1310 Cys Arg
ttg gtt gct g Leu Val Ala A 415	gcg cag gag aag Ma Gln Glu Lys	aag aag aag Lys Lys Lys 420	agc ctg ggg aag Ser Leu Gly Lys 425	cgg ctc 1358 Arg Leu
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cca gcc tcc c Pro Ala Ser L 445	ett ctt cat cca Leu Leu His Pro 450	ctc ttt cag Leu Phe Gln	gca atg act gac Ala Met Thr Asp 455	cca ctg 1454 Pro Leu 460
tgc cat ctg a Cys His Leu S	gc agc ctc acg Ser Ser Leu Thr 465	ctg tcc cac Leu Ser His 470	tgc aaa ctc cct Cys Lys Leu Pro	gac gcg 1502 Asp Ala 475
Val Cys Arg A	ac ctt tct gag sp Leu Ser Glu 80	gcc ctg agg Ala Leu Arg 485	gca gcc ccc gca Ala Ala Pro Ala 490	ctg acg 1550 Leu Thr
gag ctg ggc c Glu Leu Gly L 495	etc ctc cac aac eu Leu His Asn	agg ctc agt Arg Leu Ser 500	gag gcg gga ctg Glu Ala Gly Leu 505	cgt atg 1598 Arg Met
ctg agt gag g Leu Ser Glu G 510	gc cta gcc tgg ly Leu Ala Trp 515	ccg cag tgc Pro Gln Cys	agg gtg cag acg Arg Val Gln Thr 520	gtc agg 1646 Val Arg
Val Gln Leu P. 525	ro Asp Pro Gln 530	Arg Gly Leu	cag tac ctg gtg Gln Tyr Leu Val 535	Gly Met 540
Leu Arg Gln S	er Pro Ala Leu 545	Thr Thr Leu 550		Cys Gln 555
Leu Pro Ala P	ro Met Val Thr 60	Tyr Leu Cys 565	gca gtc ctg cag Ala Val Leu Gln 570	His Gln
Gly Cys Gly Le 575	eu Gln Thr Leu	Ser Leu Ala 580	tct gtg gag ctg Ser Val Glu Leu 585	Ser Glu
Gln Ser Leu Gl 590	ln Glu Leu Gln 595	Ala Val Lys	aga gca aag ccg Arg Ala Lys Pro . 600	Asp Leu
gtc atc aca ca Val Ile Thr Hi 605	ac cca gcg ctg is Pro Ala Leu 610	Asp Gly His	cca caa cct ccc Pro Gln Pro Pro 615	aag gaa 1934 Lys Glu 620

ctc atc to	g acc ttc to r Thr Phe 625	gaggetetg g	tggccagag ca	agggtggaa ga	accctagtc	1989
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gggaaacttt	tgagccgaga	ggccgcagac	aggcatgtgg	gaggcccaga	cacggcaccc	2109
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Pro Gly Arg Leu Gln Gly Arg Leu Cys Ser Pro Gln Cys Ala Glu Val 65 70 75 80

Arg Gly Phe Ser Asp Lys Asp Lys Lys Lys Tyr Phe Tyr Lys Phe Phe 85 90 95

Arg Asp Glu Arg Ala Glu Arg Ala Tyr Arg Phe Val Lys Glu Asn 100 105 110

Glu Thr Leu Phe Ala Leu Cys Phe Val Pro Phe Val Cys Trp Ile Val 115 120 125

Cys Thr Val Leu Arg Gln Gln Leu Glu Leu Gly Arg Asp Leu Ser Arg 130 135 140

Thr Ser Lys Thr Thr Thr Ser Val Tyr Leu Leu Phe Ile Thr Ser Val 145 150 155 160

Leu Ser Ser Ala Pro Val Ala Asp Gly Pro Arg Leu Gln Gly Asp Leu 165 170 175

Arg Asn Leu Cys Arg Leu Ala Arg Glu Gly Val Leu Gly Arg Arg Ala 180 185 190

Gln Phe Ala Glu Lys Glu Leu Glu Gln Leu Glu Leu Arg Gly Ser Lys 195 200 205

Val Gln Thr Leu Phe Leu Ser Lys Lys Glu Leu Pro Gly Val Leu Glu 210 215 220

Thr Glu Val Thr Tyr Gln Phe Ile Asp Gln Ser Phe Gln Glu Phe Leu 225 230 235 240

Ala Ala Leu Ser Tyr Leu Leu Glu Asp Gly Gly Val Pro Arg Thr Ala 245 250 255

Ala Gly Gly Val Gly Thr Leu Leu Arg Gly Asp Ala Gln Pro His Ser 260 265 270

His Leu Val Leu Thr Thr Arg Phe Leu Phe Gly Leu Leu Ser Ala Glu 275 280 285

Arg Met Arg Asp Ile Glu Arg His Phe Gly Cys Met Val Ser Glu Arg 290 295 300

Val Lys Gln Glu Ala Leu Arg Trp Val Gln Gly Gln Gly Gln Gly Cys 305 310 315 320

Pro Gly Val Ala Pro Glu Val Thr Glu Gly Ala Lys Gly Leu Glu Asp 325 330 335

Thr Glu Glu Pro Glu Glu Glu Glu Glu Glu Glu Pro Asn Tyr Pro

Leu	Glu	Leu	Leu	Tyr	Cys	Leu	Tyr	Glu	Thr	Gln	Glu	Asp	Ala	Phe	Val
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Phe Cys Arg Met Asp Val Ala Val Leu Ser Tyr Cys Val Arg Cys Cys 385 390 390 395 400

Pro Ala Gly Gln Ala Leu Arg Leu Ile Ser Cys Arg Leu Val Ala Ala 405 410 415

Gln Glu Lys Lys Lys Ser Leu Gly Lys Arg Leu Gln Ala Ser Leu 420 425 430

Gly Gly Ser Ser Gln Gly Thr Thr Lys Gln Leu Pro Ala Ser Leu 435 440 445

Leu His Pro Leu Phe Gln Ala Met Thr Asp Pro Leu Cys His Leu Ser 450 455 460

Ser Leu Thr Leu Ser His Cys Lys Leu Pro Asp Ala Val Cys Arg Asp 465 470 475 480

Leu Ser Glu Ala Leu Arg Ala Ala Pro Ala Leu Thr Glu Leu Gly Leu 485 490 495

Leu His Asn Arg Leu Ser Glu Ala Gly Leu Arg Met Leu Ser Glu Gly 500 505 510

Leu Ala Trp Pro Gln Cys Arg Val Gln Thr Val Arg Val Gln Leu Pro 515 520 525

Asp Pro Gln Arg Gly Leu Gln Tyr Leu Val Gly Met Leu Arg Gln Ser 530 540

Pro Ala Leu Thr Thr Leu Asp Leu Ser Gly Cys Gln Leu Pro Ala Pro 545 550 555 560

Met Val Thr Tyr Leu Cys Ala Val Leu Gln His Gln Gly Cys Gly Leu 565 570 575

Gln Thr Leu Ser Leu Ala Ser Val Glu Leu Ser Glu Gln Ser Leu Gln 580 585 590

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Pro Ala Leu Asp Gly His Pro Gln Pro Pro Lys Glu Leu Ile Ser Thr 610 615 620

Phe 625

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<212> PRT

<213> homo sapiens

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Lys Lys Glu Glu Leu Lys Glu Phe Gln Leu Leu Leu Ala Asn Lys Ala 20 25 30

His Ser Arg Ser Ser Ser Gly Glu Thr Pro Ala Gln Pro Glu Lys Thr 35 40 45

Ser Gly Met Glu Val Ala Ser Tyr Leu Val Ala Gln Tyr Gly Glu Gln 50 55 60

Arg Ala Trp Asp Leu Ala Leu His Thr Trp Glu Gln Met Gly Leu Arg 65 70 75 80

Ser Leu Cys Ala Gln Ala Gln Glu Gly Ala Gly His Ser Pro Ser Phe 85 90 95

Pro Tyr Ser Pro Ser Glu Pro His Leu Gly Ser Pro Ser Gln Pro Thr

Ser Thr Ala Val Leu Met Pro Trp Ile His Glu Leu Pro Ala Gly Cys 115 120 125

Thr Gln Gly Ser Glu Arg Arg Val Leu Arg Gln Leu Pro Asp Thr Ser 130 135 140

Gly Arg Arg Trp Arg Glu Ile Ser Ala Ser His Leu Tyr Gln Ala Leu 145 150 155 160

Pro Ser Ser Pro Asp His Glu Ser Pro Ser Gln Glu Ser Pro Asn Ala 165 170 175

Pro Thr Ser Thr Ala Val Leu Gly Ser Trp Gly Ser Pro Pro Gln Pro
180 185

Ser Leu Ala Pro Arg Glu Gln Glu Ala Pro Gly Thr Gln Trp Pro Leu 195 200 205

Asp Glu Thr Ser Gly Ile Tyr Tyr Thr Glu Ile Arg Glu Arg Glu Arg

Glu 225		s Ser	Glu	Lys	Gly 230		Pro	Pro	Trp	Ala 235		Val	. Val	Gly	Thi 240
Pro	Pro	Gln	Ala	His 245		Ser	Leu	Gln	Pro 250		His	His	Pro	Trp 255	
Pro	Ser	· Val	Arg 260		Ser	Leu	Cys	Ser 265		Trp	Pro	Trp	Lys 270		Glu
Asp	Phe	Asn 275	Gln	Lys	Phe	Thr	Gln 280		Leu	Leu	Leu	Gln 285		Pro	His
Pro	Arg 290	Ser	Gln	Asp	Pro	Leu 295		Lys	Arg	Ser	Trp 300		Asp	Tyr	Val
Glu 305	Glu	Asn	Arg	Gly	His 310	Leu	Ile	Glu	Ile	Arg 315	Asp	Leu	Phe	Gly	Pro 320
Gly	Leu	Asp	Thr	Gln 325	Glu	Pro	Arg	Ile	Val 330	Ile	Leu	Gln	Gly	Ala 335	Ala
Gly	Ile	Gly	Lys 340	Ser	Thr	Leu	Ala	Arg 345	Gln	Val	Lys	Glu	Ala 350	Trp	Gly
Arg	Gly	Gln 355	Leu	Tyr	Gly	Asp	Arg 360	Phe	Gln	His	Val	Phe 365	Tyr	Phe	Ser
Cys	Arg 370	Glu	Leu	Ala	Gln	Ser 375	Lys	Val	Val	Ser	Leu 380	Ala	Glu	Leu	Ile
385			Gly		390					395					400
Arg	Pro	Glu	Arg	Leu 405	Leu	Phe	Ile	Leu	Asp 410	Gly	Val	Asp	Glu	Pro 415	Gly
			Gln 420					425					430		
Pro	Gln	Pro 435	Ala	Asp	Ala	Leu	Leu 440	Gly	Ser	Leu	Leu	Gly 445	Lys	Thr	Ile
Leu	Pro 450	Glu	Ala	Ser	Phe	Leu 455	Ile	Thr	Ala	Arg	Thr 460	Thr	Ala	Leu	Gln
Asn 465	Leu	Ile	Pro	Ser	Leu 470	Glu	Gln	Ala	Arg	Trp 475	Val	Glu	Val	Leu	Gly 480
Phe	Ser	Glu	Ser	Ser 485	Arg	Lys	Glu	Tyr	Phe 490	Tyr	Arg	Tyr	Phe	Thr 495	Asp
Glu	Arg	Gln	Ala 500	Ile	Arg	Ala	Phe	Arg 505	Leu	Val	Lys	Ser	Asn 510	Lys	Glu
Leu	Trp	Ala 515	Leu	Cys	Leu	Val	Pro 520	Trp	Val	Ser	Trp	Leu 525	Ala	Cys	Thr
Cys	Leu 530	Met	Gln	Gln	Met	Lys 535	Arg	Lys	Glu	Lys	Leu 540	Thr	Leu	Thr	Ser

Lys Thr Thr Thr Leu Cys Leu His Tyr Leu Ala Gln Ala Leu Gln 545 550 560

Ala Gln Pro Leu Gly Pro Gln Leu Arg Asp Leu Cys Ser Leu Ala Ala 565 570 575

Glu Gly Ile Trp Gln Lys Lys Thr Leu Phe Ser Pro Asp Asp Leu Arg
580 585 590

Lys His Gly Leu Asp Gly Ala Ile Ile Ser Thr Phe Leu Lys Met Gly 595 600 605

Ile Leu Gln Glu His Pro Ile Pro Leu Ser Tyr Ser Phe Ile His Leu 610 615 620

Cys Phe Gln Glu Phe Phe Ala Ala Met Ser Tyr Val Leu Glu Asp Glu 625 630 635 640

Lys Gly Arg Gly Lys His Ser Asn Cys Ile Ile Asp Leu Glu Lys Thr 645 650 655

Leu Glu Ala Tyr Gly Ile His Gly Leu Phe Gly Ala Ser Thr Thr Arg 660 665 670

Phe Leu Gly Leu Leu Ser Asp Glu Gly Glu Arg Glu Met Glu Asn 675 680 685

Ile Phe His Cys Arg Leu Ser Gln Gly Arg Asn Leu Met Gln Trp Val 690 695 700

Pro Ser Leu Gln Leu Leu Gln Pro His Ser Leu Glu Ser Leu His 705 710 715 720

Cys Leu Tyr Glu Thr Arg Asn Lys Thr Phe Leu Thr Gln Val Met Ala 725 730 735

His Phe Glu Glu Met Gly Met Cys Val Glu Thr Asp Met Glu Leu Leu 740 745 750

Val Cys Thr Phe Cys Ile Lys Phe Ser Arg His Val Lys Lys Leu Gln 755 760 765

Leu Ile Glu Gly Arg Gln His Arg Ser Thr Trp Ser Pro Ser Met Val 770 780

Val Leu Phe Arg Trp Val Pro Val Thr Asp Ala Tyr Trp Gln Ile Leu 785 790 795 800

Phe Ser Val Leu Lys Val Thr Arg Asn Leu Lys Glu Leu Asp Leu Ser 805 810 815

Gly Asn Ser Leu Ser His Ser Ala Val Lys Ser Leu Cys Lys Thr Leu 820 825 830

Arg Arg Pro Arg Cys Leu Leu Glu Thr Leu Arg Leu Ala Gly Cys Gly 835 840

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- Gln Thr Leu Thr Glu Leu Asp Leu Ser Phe Asn Val Leu Met Asp Ala 865 870 875 880
- Gly Ala Lys His Leu Cys Gln Arg Leu Arg Gln Pro Ser Cys Lys Leu 885 890 895
- Gln Arg Leu Gln Leu Val Ser Cys Gly Leu Thr Ser Asp Cys Cys Gln 900 905 910
- Asp Leu Ala Ser Val Leu Ser Ala Ser Pro Ser Leu Lys Glu Leu Asp 915 920 925
- Leu Gln Gln Asn Asn Leu Asp Asp Val Gly Val Arg Leu Cys Glu 930 935 940
- Gly Leu Arg His Pro Ala Cys Lys Leu Ile Arg Leu Gly Leu Asp Gln 945 950 955 960
- Thr Thr Leu Ser Asp Glu Met Arg Gln Glu Leu Arg Ala Leu Glu Gln
 965 970 975
- Glu Lys Pro Gln Leu Leu Ile Phe Ser Arg Arg Lys Pro Ser Val Met 980 985 990
- Thr Pro Ile Glu Gly Leu Asp Thr Gly Glu Met Ser Asn Ser Thr Ser 995 1000 1005
- Ser Leu Lys Arg Gln Arg Leu Gly Ser Glu Arg Ala Ala Ser His 1010 \$1015\$
- Val Ala Gln Ala Asn Leu Lys Leu Leu Asp Val Ser Lys Ile Phe 1025 1030 1035
- Pro Ile Ala Glu Ile Ala Glu Glu Ser Ser Pro Glu Val Val Pro 1040 1045 1050
- Val Glu Leu Leu Cys Val Pro Ser Pro Ala Ser Gln Gly Asp Leu 1055 1060 1065
- His Thr Lys Pro Leu Gly Thr Asp Asp Asp Phe Trp Gly Pro Thr 1070 1075 1080
- Gly Pro Val Ala Thr Glu Val Val Asp Lys Glu Lys Asn Leu Tyr 1085 1090 1095
- Arg Val His Phe Pro Val Ala Gly Ser Tyr Arg Trp Pro Asn Thr 1100 1105 1110
- Gly Leu Cys Phe Val Val Arg Glu Ala Val Thr Val Glu Ile Glu 1115 1120 1125
- Phe Cys Val Trp Asp Gln Phe Leu Gly Glu Ile Asn Pro Gln His 1130 1135 1140
- Ser Trp Met Val Ala Gly Pro Leu Leu Asp Ile Lys Ala Glu Pro 1145 1155
- Gly Ala Val Glu Ala Val His Leu Pro His Phe Val Ala Leu Gln 1160 1165 1170
- Gly Gly His Val Asp Thr Ser Leu Phe Gln Val Ala His Phe Lys

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His	Ile 1205	Val	Leu	Glu	Asn	Pro 1210		Phe	Ser	Pro	Leu 1215	Gly	Val	Leu
Leu	Lys 1220	Met	Ile	His	Asn	Ala 1225		Arg	Phe	Ile	Pro 1230	Val	Thr	Ser
Val	Val 1235	Leu	Leu	Tyr	His	Arg 1240		His	Pro	Glu	Glu 1245	Val	Thr	Phe
His	Leu 1250	Tyr	Leu	Ile	Pro	Ser 1255		Cys	Ser	Ile	Arg 1260		Glu	Leu
Glu	Leu 1265	Cys	Tyr	Arg	Ser	Pro 1270		Glu	Asp	Gln	Leu 1275	Phe	Ser	Glu
Phe	Tyr 1280	Val	Gly	His	Leu	Gly 1285	Ser	Gly	Ile	Arg	Leu 1290	Gln	Val	Lys
Asp	Lys 1295	Lys	Asp	Glu	Thr	Leu 1300	Val	Trp	Glu	Ala	Leu 1305	Val	Lys	Pro
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Ala	Val 1325	Pro	Ser	Pro	Leu	Asp 1330	Ala	Pro	Gln	Leu	Leu 1335	His	Phe	Val
Asp	Gln 1340	Tyr	Arg	Glu	Gln	Leu 1345	Ile	Ala	Arg	Val	Thr 1350	Ser	Val	Glu
Val	Val 1355	Leu	Asp	Lys	Leu	His 1360	Gly	Gln	Val	Leu	Ser 1365	Gln	Glu	Gln
Tyr	Glu 1370	Arg	Val	Leu	Ala	Glu 1375	Asn	Thr	Arg	Pro	Ser 1380	Gln	Met	Arg
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Gly	Leu 1400	Tyr	Gln	Ala	Leu	Lys 1405	Glu	Thr	His	Pro	His 1410	Leu	Ile	Met
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Leu	Val 50	Glu	Ile	Leu	Thr	Thr 55	His	Cys	Asp	Ser	Tyr 60	Trp	Val	Glu	Met
Ala 65	Ser	Leu	Gln	Val	Phe 70	Glu	Lys	Met	His	Arg 75	Met	Asp	Leu	Ser	Glu 80
Arg	Ala	Lys	Asp	Glu 85	Val	Arg	Glu	Ala	Ala 90	Leu	Lys	Ser	Phe	Asn 95	Lys
Arg	Lys	Pro	Leu 100	Ser	Leu	Gly	Ile	Thr 105	Arg	Lys	Glu	Arg	Pro 110	Pro	Leu
Asp	Val	Asp 115	Glu	Met	Leu	Glu	Arg 120	Phe	Lys	Thr	Glu	Ala 125	Gln	Asp	Lys
Asp	Asn 130	Arg	Cys	Arg	Tyr	Ile 135	Leu	Lys	Thr	Lys	Phe 140	Arg	Glu	Met	Trp
Lys 145	Ser	Trp	Pro	Gly	Asp 150	Ser	Lys	Glu	Val	Gln 155	Val	Met	Ala	Glu	Arg 160
Tyr	Lys	Met	Leu	Ile 165	Pro	Phe	Ser	Asn	Pro 170	Arg	Val	Leu	Pro	Gly 175	Pro
Phe	Ser	Tyr	Thr 180	Val	Val	Leu	Tyr	Gly 185	Pro	Ala	Gly	Leu	Gly 190	Lys	Thr
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Leu 225	Gly	Pro	Cys	Ser	Phe 230	Ala	Glu	Leu	Val	Phe 235	Arg	Asp	Trp	Pro	Glu 240
	Gln			245					250					255	
Phe	Val	Ile	Asp 260	Gly	Phe	Asp	Glu	Leu 265	Gly	Ala	Ala	Pro	Gly 270	Ala	Leu
Ile	Glu	Asp 275	Ile	Cys	Gly	Asp	Trp 280	Glu	Lys	Lys	Lys	Pro 285	Val	Pro	Val
Leu	Leu 290	Gly	Ser	Leu	Leu	Asn 295	Arg	Val	Met	Leu	Pro 300	Lys	Ala	Ala	Leu
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Ser Leu Cys Ala Gln Ala Gln Glu Gly Ala Gly His Ser Pro Ser Phe 85 90 95

Pro Tyr Ser Pro Ser Glu Pro His Leu Gly Ser Pro Ser Gln Pro Thr 100 105 110

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Thr Gln Gly Ser Glu Arg Arg Val Leu Arg Gln Leu Pro Asp Thr Ser 130 135 140

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Glv	T. 211	Δνα	шic	Dwo	חות	Q	т	_	~ 7	_	_				

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